



1/2

SEQUENCE LISTING

COPY OF PAPERS
ORIGINALY FILED

<110> Aevarsson, Arnthor
Marteinsson, Viggo T.
Hreggvidsson, Gudmundur O.
Kristjansson, Jakob K.
Fridjonsson, Olafur H.

<120> Method of Obtaining Protein Diversity

<130> 2739.2004-001

<140> 09/878,423

<141> 2001-06-11

<150> Iceland 5863

<151> 2001-02-23

<160> 6

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> bacterial primer

<400> 1

gagtttgatc ctggctcag

19

<210> 2

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> bacterial primer

<400> 2

gtcccagcag ccgcggtaaa tac

23

<210> 3

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> bacterial primer

<400> 3

gactaccggg tatctaacc

20

<210> 4
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> bacterial primer

<400> 4
agaaaggagg tgatcca

17

<210> 5
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> archaea primer

<400> 5
gcggatccgc ggccgctgca gayctggtyg atyctgcc

38

<210> 6
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> archaea primer

<400> 6
gacgggagggt gtgtrca

17